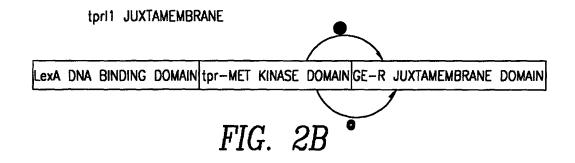
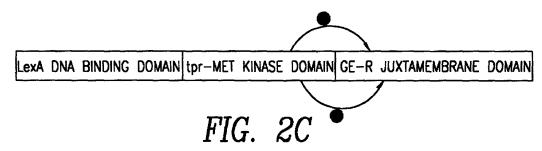


FIG. 2A



tprl1 C-TERMINAL



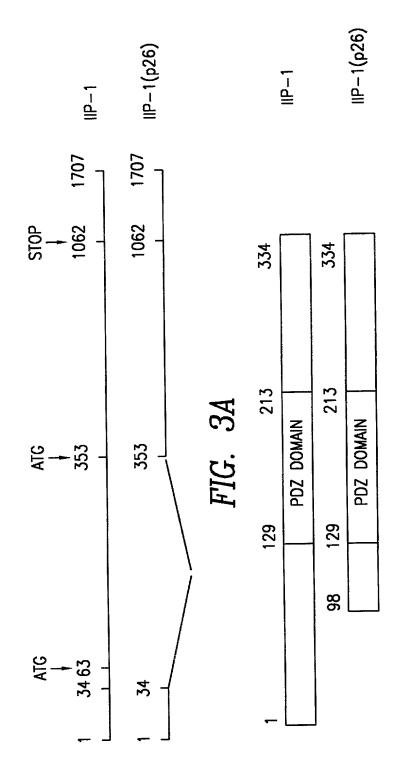


FIG. 3B

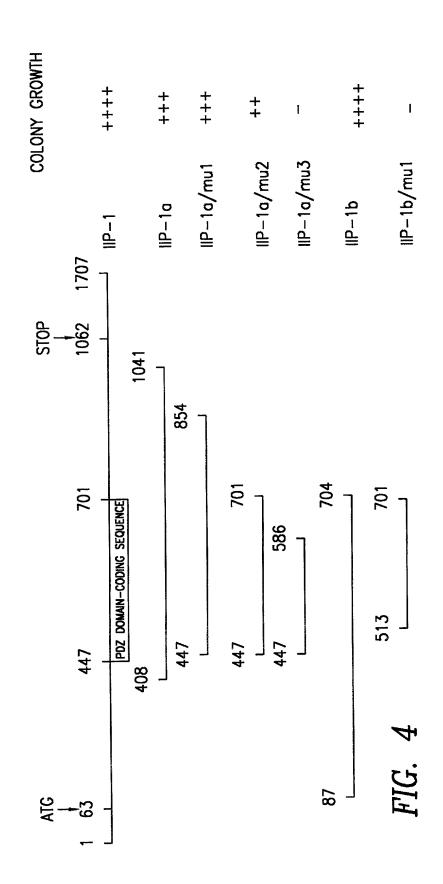


Fig. 5

- 1 MSRPRKRLAG TSGSBKGLSG KRTKTENSGE ALAKVEDSNP QKTSATKNCL
- 51 KNLSSHWLMK SEPESRLEKG VDVKFSTEDL KAQPKQTTCW DGVRNYQARN
- ↑
 101 FLRAMKLGEE AFFYHSNCKE PGIAGLMKIV KEAYPDHTOF EKNNPHYDPS
- 151 SKEDNPKWSM VDVQFVRMMK RFIPLAELKS YHQAHKATGG PLKNMVLFTR
- 201 QRLSIQPLTQ EEFDFVLSTE EKEPS*
- putative N-glycosylation site
- putative Casein kinase II phosphorylation site
- putative N-myristoylation site
- putative Tyrosine kinase phosphorylation site
- putative Protein kinase C phosphorylation site
- ←→ putative NLS

